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SEQUENCE LISTING

TECH CENTER 1600 2300

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<110> MAYAGAWA, SHUJI MADEMAN KABE, MASARU

<120> MODIFIED CRE RECOMBINASE GENE FOR MAMMALS
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<141> 2000-09-14
<150> JP11-264364
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<170> PatentIn version 3.0

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tac ctg ctg tac ctg cag gcc cgc ggc ctg gcc gtg aag acc atc cag

Tyr Leu Leu Tyr Leu Gln Ala Arg Gly Leu Ala Val Lys Thr Ile Gln

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ggc tgg acc aac gtg aac atc gtg atg aac tac atc cgc aac ctg gac Gly Trp Thr Asn Val Asn Ile Val Met Asn Tyr Ile Arg Asn Leu Asp 325 age gag ace gge gee atg gtg ege etg etg gag gae gge gae Ser Glu Thr Gly Ala Met Val Arg Leu Leu Glu Asp Gly Asp 340 345 <210> 2 <211> 350 <212> PRT <213> Artificial Sequence <220> <221> misc_feature <222> ()..() <223> Description of Artificial Sequence: synthetic peptide <400> 2 Met Pro Lys Lys Lys Arg Lys Val Ser Asn Leu Leu Thr Val His Gln 5 10 Asn Leu Pro Ala Leu Pro Val Asp Ala Thr Ser Asp Glu Val Arg Lys 25 Asn Leu Met Asp Met Phe Arg Asp Arg Gln Ala Phe Ser Glu His Thr 40 Trp Lys Met Leu Leu Ser Val Cys Arg Ser Trp Ala Ala Trp Cys Lys 50 Leu Asn Asn Arg Lys Trp Phe Pro Ala Glu Pro Glu Asp Val Arg Asp Tyr Leu Leu Tyr Leu Gln Ala Arg Gly Leu Ala Val Lys Thr Ile Gln Gln His Leu Gly Gln Leu Asn Met Leu His Arg Arg Ser Gly Leu Pro 100 Arg Pro Ser Asp Ser Asn Ala Val Ser Leu Val Met Arg Arg Ile Arg 115 120 Lys Glu Asn Val Asp Ala Gly Glu Arg Ala Lys Gln Ala Leu Ala Phe

1008

1050

135

130

Glu Arg Thr Asp Phe Asp Gln Val Arg Ser Leu Met Glu Asn Ser Asp 145 150 155 160

Arg Cys Gln Asp Ile Arg Asn Leu Ala Phe Leu Gly Ile Ala Tyr Asn 165 170 175

Thr Leu Leu Arg Ile Ala Glu Ile Ala Arg Ile Arg Val Lys Asp Ile 180 185 190

Ser Arg Thr Asp Gly Gly Arg Met Leu Ile His Ile Gly Arg Thr Lys
195 200 205

Thr Leu Val Ser Thr Ala Gly Val Glu Lys Ala Leu Ser Leu Gly Val 210 220

Thr Lys Leu Val Glu Arg Trp Ile Ser Val Ser Gly Val Ala Asp Asp 225 230 230 235

Pro Asn Asn Tyr Leu Phe Cys Arg Val Arg Lys Asn Gly Val Ala Ala 245 250 255

Pro Ser Ala Thr Ser Gln Leu Ser Thr Arg Ala Leu Glu Gly Ile Phe 260 265 270

Glu Ala Thr His Arg Leu Ile Tyr Gly Ala Lys Asp Asp Ser Gly Gln 275 280 285

Arg Tyr Leu Ala Trp Ser Gly His Ser Ala Arg Val Gly Ala Ala Arg 290 295 300

Asp Met Ala Arg Ala Gly Val Ser Ile Pro Glu Ile Met Gln Ala Gly 305 310 315 320

Gly Trp Thr Asn Val Asn Ile Val Met Asn Tyr Ile Arg Asn Leu Asp 325 330 335

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